## SEQUENCE LIST AP20 Rec'd PCT/PTO 12 JUN 2006

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<110> Berzofsky, Jay A.
      Pastan, Ira H.
      Terabe, Masaki
      The Government of the United States of America
         as represented by The Secretary of the
         Department of Health and Human Services
<120> Immunogenic Peptides of XAGE-1
<130> 015280-485100PC
<140> WO PCT/US04/41639
<141> 2004-12-13
<150> US 60/529,025
<151> 2003-12-12
<160> 45
<170> PatentIn Ver. 2.1
<210> 1
<211> 246
<212> DNA
<213> Homo sapiens
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<223> xage-1 p9, 9kD protein expressed from XAGE-1 gene
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<222> (1)..(246)
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atg gag agc ccc aaa aag aag aac cag cag ctg aaa gtc ggg atc cta
                                                                    48
Met Glu Ser Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu
 1
                                                          15
cac ctg ggc agc aga cag aag atc agg ata cag ctg aga tcc cag
His Leu Gly Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln
             20
                                 25
                                                      3.0
tgc gcg aca tgg aag gtg atc tgc aag agc tgc atc agt caa aca ccg
Cys Ala Thr Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro
         35
                             40
ggg ata aat ctg gat ttg ggt tcc ggc gtc aag gtg aag ata ata cct
Gly Ile Asn Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro
     50
                         55
aaa gag gaa cac tgt aaa atg cca gaa gca ggt gaa gag caa cca caa
Lys Glu Glu His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln
65
                     70
                                          75
gtt taa
                                                                   246
Val
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<210> 2
<211> 81
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<213> Homo sapiens
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<400> 2
Met Glu Ser Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu
His Leu Gly Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln
Cys Ala Thr Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro
Gly Ile Asn Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro
Lys Glu Glu His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln
Val
<210> 3
<211> 441
<212> DNA
<213> Homo sapiens
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<223> xage-1 p16, 16.3 kD protein expressed from XAGE-1
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<222> (1) .. (441)
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<400> 3
atg etc ett tgg tge eea eet eag tge gea tgt tea etg gge gte tte
                                                                   48
Met Leu Leu Trp Cys Pro Pro Gln Cys Ala Cys Ser Leu Gly Val Phe
cca tcg gcc cct tcg cca gtg tgg gga acg cgg cgg agc tgt gag ccg
                                                                   96
Pro Ser Ala Pro Ser Pro Val Trp Gly Thr Arg Arg Ser Cys Glu Pro
gcg act cgg gtc cct gag gtc tgg att ctt tct ccg cta ctg aga cac
                                                                   144
Ala Thr Arg Val Pro Glu Val Trp Ile Leu Ser Pro Leu Leu Arg His
                             40
ggc gga cac aca caa aca cag aac cac aca gcc agt ccc agg agc cca
                                                                   192
Gly Gly His Thr Gln Thr Gln Asn His Thr Ala Ser Pro Arg Ser Pro
gta atg gag agc ccc aaa aag aag aac cag cag ctg aaa gtc ggg atc
Val Met Glu Ser Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile
65
                     70
                                          75
```

cta Leu	cac His	ctg Leu	ggc Gly	agc Ser 85	Arg	cag Gln	aag Lys	aag Lys	atc Ile 90	agg Arg	ata Ile	cag Gln	ctg Leu	aga Arg 95	tcc Ser	288
cag Gln	tgc Cys	gcg Ala	aca Thr 100	tgg Trp	aag Lys	gtg Val	atc Ile	tgc Cys 105	aag Lys	agc Ser	tgc Cys	atc Ile	agt Ser 110	caa Gln	aca Thr	336
	gj aaa															384
cct Pro	aaa Lys 130	gag Glu	gaa Glu	cac His	tgt Cys	aaa Lys 135	atg Met	cca Pro	gaa Glu	gca Ala	ggt Gly 140	gaa Glu	gag Glu	caa Gln	cca Pro	432
	gtt Val	taa														441
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Pro	Ser	Ala	Pro 20	Ser	Pro	Val	Trp	Gly 25	Thr	Arg	Arg	Ser	Cys 30	Glu	Pro	
Ala	Thr	Arg 35	Val	Pro	Glu	Val	Trp 40	Ile	Leu	Ser	Pro	Leu 45	Leu	Arg	His	
Gly	Gly 50	His	Thr	Gln	Thr	Gln 55	Asn	His	Thr	Ala	Ser 60	Pro	Arg	Ser	Pro	
Val 65	Met	Glu	Ser	Pro	Lys 70	Lys	Lys	Asn	Gln	Gln 75	Leu	Lys	Val	Gly	Ile 80	
Leu	His	Leu	Gly	Ser 85	Arg	Gln	Lys	Lys	Ile 90	Arg	Ile	Gln	Leu	Arg 95	Ser	
				05												
Gln	Cys	Ala	Thr 100		Lys	Val	Ile	Cys 105	Lys	Ser	Сув	Ile	Ser 110	Gln	Thr	
	Cys Gly		100	Trp				105					110			
Pro		Ile 115	100 Asn	Trp Leu	Asp	Leu	Gly 120	105 Ser	Gly	Val	Lys	Val 125	110 Lys	Ile	Ile	

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<210> 5
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<212> PRT
<213> Artificial Sequence
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<221> MOD RES
<222> (1)
<223> Xaa = any amino acid (X-1)
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<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
<220>
<221> MOD RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD_RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 5
Xaa Xaa Xaa Pro Ser Ala Pro Ser Pro Xaa
                  5
<210> 6
<211> 10
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:xage-1 14,
      immunogenic amino terminal end of xage-1, xage-1
      residues 14-23
<400> 6
Gly Val Phe Pro Ser Ala Pro Ser Pro Val
<210> 7
<211> 10
<212> PRT
<213> Artificial Sequence
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      variant of xage-1 14, immunogenic peptide derived
      from xage-1 14
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<400> 7
Tyr Val Phe Pro Ser Ala Pro Ser Pro Val
<210> 8
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: 2L xage-1 14,
      variant of xage-1 14, immunogenic peptide derived
      from xage-1 14
<400> 8
Gly Leu Phe Pro Ser Ala Pro Ser Pro Val
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<210> 9
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:3M xage-1 14,
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<400> 9
Gly Val Met Pro Ser Ala Pro Ser Pro Val
<210> 10
<211> 10
<212> PRT
<213> Artificial Sequence
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      from xage-1 14
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Tyr Leu Phe Pro Ser Ala Pro Ser Pro Val
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                  5
<210> 11
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<212> PRT
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<220>
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      variant of xage-1 14, immunogenic peptide derived
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Gly Leu Met Pro Ser Ala Pro Ser Pro Val
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<210> 12
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<223> Description of Artificial Sequence:modified xage-1
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Gly Val Trp Pro Ser Ala Pro Ser Pro Val
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<210> 13
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: modified xage-1
      14 peptide, immunogenic peptide derived from
      xage-1 14
<400> 13
Gly Val Tyr Pro Ser Ala Pro Ser Pro Val
<210> 14
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     xage-1 14
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Thr Val Trp Pro Ser Ala Pro Ser Pro Met
<210> 15
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     xage-1 14
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Ser Met Tyr Pro Ser Ala Pro Ser Pro Ile
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<210> 16
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Ser Val Phe Pro Ser Ala Pro Ser Pro Thr
<210> 17
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Gly Val Trp Pro Ser Ala Pro Ser Pro Met
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<400> 18
Ser Val Trp Pro Ser Ala Pro Ser Pro Val
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<210> 19
<211> 10
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:modified xage-1
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     xage-1 14
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Gly Leu Trp Pro Ser Ala Pro Ser Pro Val
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<210> 20
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<210> 21
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<210> 22
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<210> 23
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      xage-1 14
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Tyr Leu Phe Pro Ser Ala Pro Ser Pro Met
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Tyr Leu Ala Pro Ser Ala Pro Ser Pro Ile
<210> 25
<211> 10
<212> PRT
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<400> 25
Tyr Leu Ala Pro Ser Ala Pro Ser Pro Val
                  5
<210> 26
<211> 30
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: nucleic acid
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ggcgtcttcc catcggcccc ttcgccagtg
                                                                    30
<210> 27
<211> 30
<212> DNA
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<220>
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<400> 27
ggcgtcatgc catcggcccc ttcgccagtg
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<210> 28
<211> 30
<212> DNA
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      sequence encoding SEQ ID NO:11 preferred form
ggccttatgc catcggcccc ttcgccagtg
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<210> 29
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: nucleic acid
      sequence encoding SEQ ID NO:11 preferred form
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ggcctcatgc catcggcccc ttcgccagtq
                                                                    30
<210> 30
<211> 30
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence:nucleic acid
      sequence encoding SEQ ID NO:11 preferred form
ggcctaatgc catcggcccc ttcgccagtg
                                                                    30
<210> 31
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
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      sequence encoding SEQ ID NO:11 preferred form
<400> 31
ggcctgatgc catcggcccc ttcgccagtg
                                                                    30
<210> 32
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<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:xage-1 33,
      residues 33-42 of xage-1
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<400> 32
Ala Thr Arg Val Pro Glu Val Trp Ile Leu
                  5
<210> 33
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:xage-1 57,
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His Thr Ala Ser Pro Arg Ser Pro Val Met
<210> 34
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<222> (2)
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<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
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Tyr Xaa Xaa Pro Ser Ala Pro Ser Pro Xaa
 1
                  5
<210> 35
<211> 10
<212> PRT
<213> Artificial Sequence
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<220>
<221> MOD RES
<222> (1)
<223> Xaa = any amino acid (X-1)
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<220>
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<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD_RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 35
Xaa Leu Xaa Pro Ser Ala Pro Ser Pro Xaa
<210> 36
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<212> PRT
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<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
<221> MOD RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 36
Xaa Xaa Met Pro Ser Ala Pro Ser Pro Xaa
<210> 37
<211> 10
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:immunogenic
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<221> MOD RES
<222> (1)
<223> Xaa = any amino acid (X-1)
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<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
<220>
<221> MOD_RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
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<210> 38
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:9-mer created
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<222> (1)
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<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
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<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD RES
<222> (9)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 38
Xaa Xaa Xaa Pro Ser Ala Pro Ser Xaa
<210> 39
<211> 9
<212> PRT
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<223> Description of Artificial Sequence:9-mer created
      from SEQ ID NO:5 by omitting Ser at position 8
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<222> (1)
<223> Xaa = any amino acid (X-1)
<220>
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<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
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<221> MOD RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
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<221> MOD_RES
<222> (9)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 39
Xaa Xaa Xaa Pro Ser Ala Pro Pro Xaa
<210> 40
<211> 9
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<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
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<221> MOD_RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD RES
<222> (9)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 40
Xaa Xaa Xaa Pro Ser Ala Ser Pro Xaa
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<210> 41
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:overall formula
      for 9-mers created from SEQ ID NO:5
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<222> (1)
<223> Xaa = any amino acid (X-1)
<220>
<221> MOD_RES
<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
<220>
<221> MOD_RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD RES
<222> (7)
<223> Xaa = Pro or absent (X-5), when absent, X-6 is Ser
<220>
<221> MOD RES
<222> (8)
<223> Xaa = Ser or absent (X-6), when absent, X-5 and X-7 are Pro
<220>
<221> MOD_RES
<222> (9)
<223> Xaa = Pro or absent (X-7), when absent, X-5 is Pro and X-6
      is Ser
<220>
<221> MOD RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 41
Xaa Xaa Xaa Pro Ser Ala Xaa Xaa Xaa
                 5
<210> 42
<211> 27
<212> DNA
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<220>
<223> Description of Artificial Sequence: exemplar
      nucleic acid encoding a peptide of SEQ ID NO:39
<400> 42
ggcgtcttcc catcggcccc ttcggtg
                                                                   27
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<210> 43
<211> 27
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: exemplar
      nucleic acid encoding a peptide of SEQ ID NO:38
<400> 43
ggcgtcttcc catcggcccc tccagtg
                                                                   27
<210> 44
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:exemplar
      nucleic acid encoding a peptide of SEQ ID NO:40
<400> 44
ggcgtcttcc catcggcctc gccagtg
                                                                   27
<210> 45
<211> 637
<212> DNA
<213> Homo sapiens
<220>
<223> complete nucleic acid sequence of XAGE-1 with
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gtcgttaatg gggacctggg aaggagcata ggacagggca aggcgggata aggagggca 60
ccacagccct taaggcacga gggaacctca ctgcgcatgc tcctttggtg cccacctcag 120
tgegcatgtt cactgggcgt cttcccateg gccccttcgc cagtgtgggg aacgcggcgg 180
agetgtgage eggegaeteg ggteeetgag gtetggatte ttteteeget aetgagaeae 240
ggeggacaca cacaaacaca gaaccacaca gccagtccca ggagcccagt aatggagagc 300
cccaaaaaga agaaccagca gctgaaagtc gggatcctac acctgggcag cagacagaag 360
aagatcagga tacagctgag atcccagtgc gcgacatgga aggtgatctg caagagctgc 420
atcagtcaaa caccggggat aaatctggat ttgggttccg gcgtcaaggt gaagataata 480
cctaaagagg aacactgtaa aatgccagaa gcaggtgaag agcaaccaca agtttaaatg 540
aagacaagct gaaacaacgc aagctggttt tatattagat atttgactta aactatctca 600
ataaagtttt gcagctttca ccaaaaaaa aaaaaaa
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